Fig. 1A

ATGAAGCTCGCCGCCCTCCTGGGGCTCTGCGTGGCCCTGTCCTGCAGCTCCGC
TCGTGCTTTCTTAGTGGGCTCGGCCAAGCCTGTGGCCCAGCCTGTCGCTGCGC
TGGAGTCGGCGGCGGAGGCCGGGGCCGGGACCCTGGCCAACCCCCTCGGCA
CCCTCAACCCGCTGAAGCTCCTGCTGAGCAGCCTGGGCATCCCCGTGAACCA
CCTCATAGAGGGCTCCCAGAAGTGTGTGGCTGAGCTGGGTCCCCAGGCCGTG
GGGGCCGTGAAGGCCCTGAAGGCCCTGCTGGGGGCCCTGACAGTGTTTGGC

Fig. 1B

CGTGCTTTCTTAGTGGGCTCGGCCAAGCCTGTGGCCCAGCCTGTCGCTGCGCTGGAGTCGGCGGGAGGCCGGGGCCGGGACCCTGGCCAACCCCCTCGGCACCCTCAACCCGCTGAAGCTCCTGCTGAGCAGCCTGGGCATCCCCGTGAACCACCTCATAGAGGGCTCCCAGAAGTGTGTGGCTGAGCTGGGTCCCCAGGCCGTGGGGCCCTGAAGGCCCTGAAGGCCCTGACAGTGTTTGGC

Fig. 1C

TTCTTAGTGGGCTCGGCCAAGCCTGTGGCCCAGCCTGTCGCTGCGCTGAGTC GGCGGCGGAGGCCGGGGCCGGGACCCTGGCCAACCCCCTCGGCACCCTCAAC CCGCTGAAGCTCCTGCTGAGCAGCCTGGGCATCCCCGTGAACCACCTCATAG AGGGCTCCCAGAAGTGTGTGGCTGAGCTGGGTCCCCAGGCCGTGGGGGCCGT GAAGGCCCTGAAGGCCCTGCTGGGGGCCCTGACAGTGTTTGGC

 $\label{eq:mklaallglcvalscssaraflvgsakpvaqpvaalesaaeagagtlanplgtl\\ nplklllsslgipvnhliegsqkcvaelgpqavgavkalkallgaltvfg$

Fig. 2B

RAFLVGSAKPVAQPVAALESAAEAGAGTLANPLGTLNPLKLLLSSLGIPVNHLIE GSQKCVAELGPQAVGAVKALKALLGALTVFG

Fig. 2C

FLVGSAKPVAQPVAALESAAEAGAGTLANPLGTLNPLKLLLSSLGIPVNHLIEGS QKCVAELGPQAVGAVKALKALLGALTVFG

ATGAAGCTTACCACCACCTTTCTAGTGCTCTGTGTGGCTCTGCTCAGTGACTC
TGGTGTTGCTTTCTTCATGGACTCATTGGCCAAGCCTGCGGTAGAACCCGTGG
CCGCCCTTGCTCCAGCTGCAGAGGCTGTGGCAGGGGCTGTGCCTAGCCTACC
ATTAAGCCACTTGGCCATCCTGAGGTTCATCCTGGCCAGCATGGGCATCCCAT
TGGATCCTCTCATAGAGGGATCCAGGAAGTGTGTCACCGAGCTGGGCCCTGA
GGCTGTAGGAGCTGTGAAGTCACTGCTGGGGGTCCTGACAATGTTCGGT

Fig. 3B

GTTGCTTTCTTCATGGACTCATTGGCCAAGCCTGCGGTAGAACCCGTGGCCGC CCTTGCTCCAGCTGCAGAGGCTGTGGCAGGGGCTGTGCCTAGCCTACCATTA AGCCACTTGGCCATCCTGAGGTTCATCCTGGCCAGCATGGGCATCCCATTGG ATCCTCTCATAGAGGGATCCAGGAAGTGTGTCACCGAGCTGGGCCCTGAGGC TGTAGGAGCTGTGAAGTCACTGCTGGGGGTCCTGACAATGTTCGGT

Fig. 3C

TTCTTCATGGACTCATTGGCCAAGCCTGCGGTAGAACCCGTGGCCGCCCTTGC
TCCAGCTGCAGAGGCTGTGGCAGGGGCTGTGCCTAGCCTACCATTAAGCCAC
TTGGCCATCCTGAGGTTCATCCTGGCCAGCATGGGCATCCCATTGGATCCTCT
CATAGAGGGATCCAGGAAGTGTGTCACCGAGCTGGGCCCTGAGGCTGTAGGA
GCTGTGAAGTCACTGCTGGGGGTCCTGACAATGTTCGGT

 ${\tt MKLTTTFLVLCVALLSDSGVAFFMDSLAKPAVEPVAALAPAAEAVAGAVPSLPLSHLAILRFILASMGIPLDPLIEGSRKCVTELGPEAVGAVKSLLGVLTMFG}$

Fig. 4B

VAFFMDSLAKPAVEPVAALAPAAEAVAGAVPSLPLSHLAILRFILASMGIPLDPLI EGSRKCVTELGPEAVGAVKSLLGVLTMFG

Fig. 4C

FFMDSLAKPAVEPVAALAPAAEAVAGAVPSLPLSHLAILRFILASMGIPLDPLIEG SRKCVTELGPEAVGAVKSLLGVLTMFG



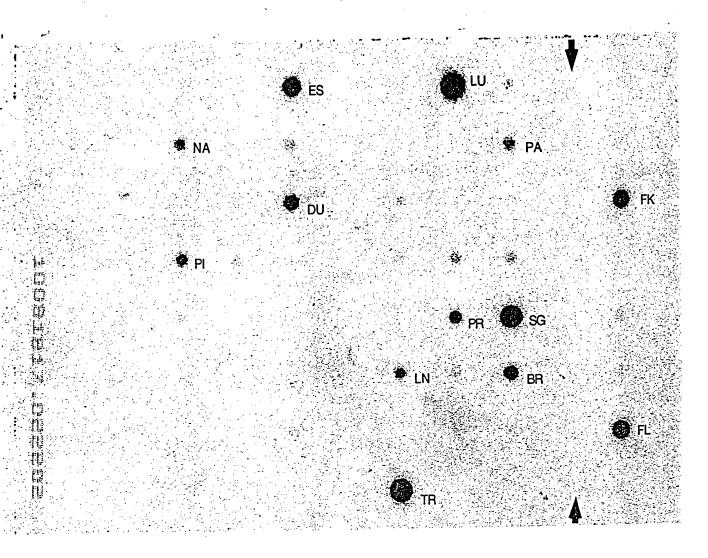


Fig. 5A

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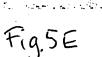
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Fig 5C



Fig.5D



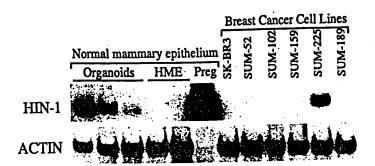
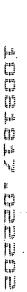


Fig. SF



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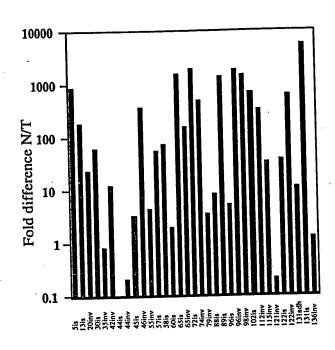
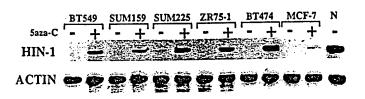


Fig. 5G.

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SK-BR-3		000000000000000000000000000000000000000		000000000	1
SUM1S9	000000000000000000000000000000000000000	200000000000000000000000000000000000000	000000000000000000000000000000000000000	000000000	1
SUM225		000000000000000000000000000000000000000		00000000	‡
T44		000000000000000000000000000000000000000	000000000000000000000000000000000000000	000000000	+
Noma		000000000000000000000000000000000000000	000000000000000000000000000000000000000	000000000	
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Fig. 7

Fig. 9A

Fig. 9B

VLCFVLVGVAFLVDSLAKPVVEPVAAIATAAEAVAGAVPSLPLSHLAILRFIVTSL GIPLDPLIDGSRKCVTELGPEAVGAVKSLLGALTTFG

Fig. 9C

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11,

TTCTTGGTGGATTCACTGGCCAAGCCTGTGGTAGAACCCGTGGCTGCCATTGC TACAGCTGCAGAGGCTGTGGCAGGGGCTGTGCCTAGCCTACCATTAAGCCAC TTGGCCATCCTGAGGTTCATCGTGACCAGCCTGGGCATCCCATTGGATCCTC CATAGATGGTTCCAGGAAGTGCGTCACCGAGCTGGGCCCTGAGGCTGTAGGA GCTGTGAAGTCACTGCTGGGGGCCCTGACAACGTTCGGT

Fig. 9D

FLVDSLAKPVVEPVAAIATAAEAVAGAVPSLPLSHLAILRFIVTSLGIPLDPLIDGS RKCVTELGPEAVGAVKSLLGALTTFG

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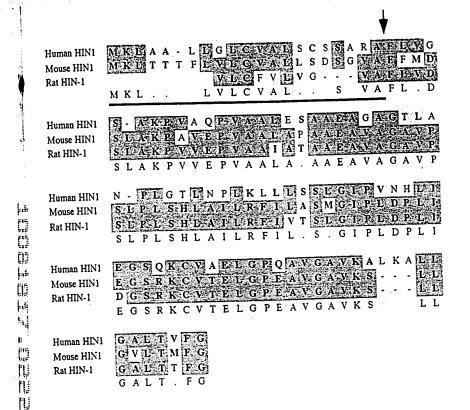


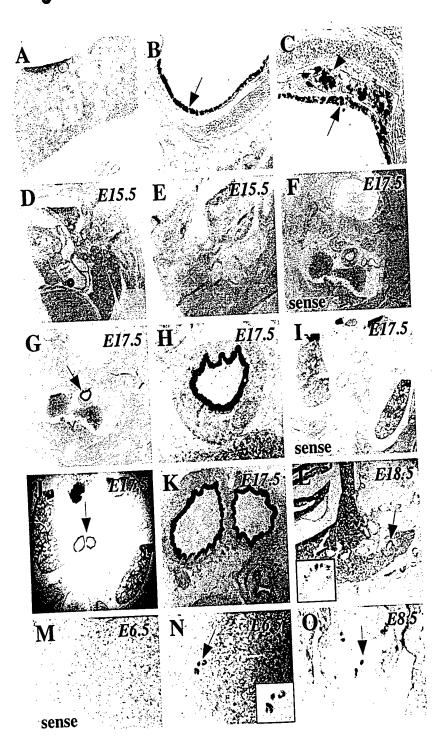
Fig. 10

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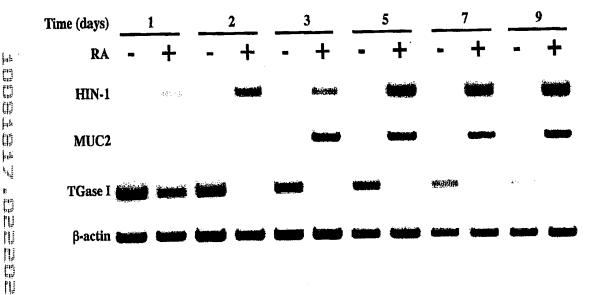
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Fig. 12



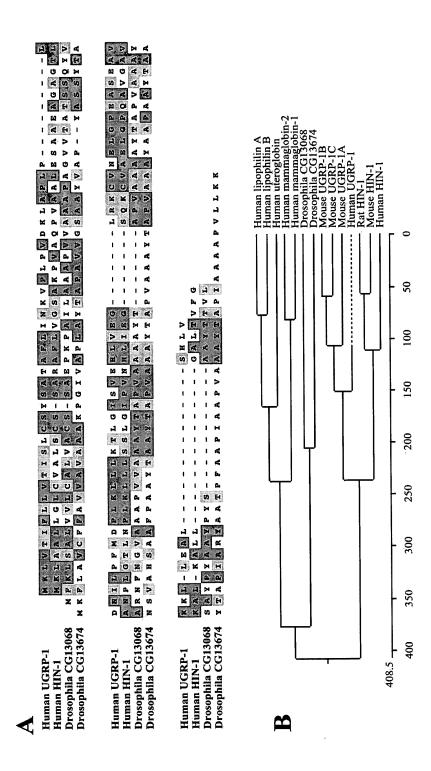
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Fig. 14

Fig. 15



Fig. 16

ATGAAATTCCTCGCCGTCTGCTTCTTCGCTGTTGTGGCTGCTGCCAA ACCCGGTATTGTGGCTCCTCTGGCCTACACCGCTCCGGCTGTGGTGGCCAGTG CCGCCTACGTGGCTCCCTACGCCTCCAGCTACACCGCCAACTCGGTGGCCCAC AGCGCCGCCTTCCCAGCTGCCTACACCGCCGCCTACACTGCTCCCGTTGCTGC TGCCTATACCGCTCCAGTGGCTGCTTATACCGCTCCAGTGGCCGCTGCGT ACGCCGCCCCAGCTGCCTATACCGCTGCCTACACCGCCCCCATTGCCCGTTAT GCCGCCACCCCCTTCGCAGCACCCATCGCCGCTCCCGTGGCTGCCGCCTACAC CGCCCCCATCGCCGCCGCTGCCCCAGTTCTGCTGAAGAAG